

1	234	100.0	1386	4	US-08-897-340-1	Sequence 1, Appl
2	227.8	97.4	871	2	US-08-744-026-2	Sequence 2, Appl
3	227.8	97.4	871	4	US-09-102-732-2	Sequence 2, Appl
4	227.8	97.4	871	6	US-08-824-874-2	Sequence 2, Appl
5	37.4	16.0	833	3	US-08-790-137-2	Sequence 2, Appl
6	37.4	16.0	833	5	US-09-025-059-2	Sequence 2, Appl
7	35.8	15.3	957	2	US-08-684-862-11	Sequence 11, Appl
8	34.8	14.9	840	2	US-08-684-862-12	Sequence 12, Appl
9	34.4	14.7	1454	2	US-08-467-155A-2	Sequence 2, Appl
10	34.4	14.7	1454	6	US-08-628-198-2	Sequence 2, Appl
11	34.4	14.7	1454	5	PCT-US96-07343-2	Sequence 9, Appl
12	34.2	14.6	1353	2	US-08-664-862-9	Sequence 9, Appl
13	33.2	14.2	1096	2	US-08-664-862-8	Sequence 8, Appl
14	32.6	13.9	986	3	US-08-557-146-1	Sequence 1, Appl
15	32.6	13.9	986	4	US-09-154-344-1	Sequence 1, Appl
16	32.6	13.9	1089	5	US-08-930-188-1	Sequence 1, Appl
17	32.6	13.9	1089	5	US-08-930-188-3	Sequence 1, Appl
18	32.6	13.9	1089	6	PCT-US96-04294-1	Sequence 3, Appl
19	32.6	13.9	1089	6	PCT-US96-04294-3	Sequence 3, Appl
20	31.6	13.5	732	2	US-08-361-395-2	Sequence 10, Appl
21	31.6	13.5	988	2	US-08-684-862-10	Sequence 10, Appl
22	30.2	12.9	1243	3	US-08-656-177A-1	Sequence 1, Appl
23	30.2	12.8	1719	3	US-08-844-024-1	Sequence 1, Appl
24	30.2	12.8	1729	4	US-08-718-547-1	Sequence 1, Appl
25	30.2	12.8	3224	5	US-09-079-415-3	Sequence 3, Appl
26	29.8	12.7	2830	1	US-07-862-792-1	Sequence 1, Appl

C 27	29.8	12.7	2830	4	US-08-331-644-1	Sequence 1, Appl1
C 28	29.8	12.7	2830	6	PCT-US93-04102-1	Sequence 1, Appl1
C 29	29	12.4	1190	1	US-08-310-370-1	Sequence 1, Appl1
C 30	28.4	12.1	992	1	US-08-358-782D-13	Sequence 13, Appl1
C 31	28.4	12.1	992	4	US-08-764-527A-13	Sequence 13, Appl1
C 32	28.4	12.1	1462	1	US-08-358-782D-14	Sequence 14, Appl1
C 33	28.4	12.1	1462	4	US-08-764-527B-14	Sequence 14, Appl1
C 34	28	12.0	2483	3	US-08-177-109A-1	Sequence 1, Appl1
C 35	28	12.0	2483	4	US-08-687-706-1	Sequence 1, Appl1
C 36	28	12.0	2634	5	US-08-941-936-1	Sequence 1, Appl1
C 37	27.8	11.9	665	4	US-08-980-060-12	Sequence 12, Appl1
C 38	27.8	11.9	3090	2	US-08-556-291-2	Sequence 2, Appl1
C 39	27.8	11.9	3090	5	US-09-100-804-2	Sequence 2, Appl1
C 40	27.8	11.9	3090	6	PCT-US94-05949-3	Sequence 3, Appl1
C 41	27.8	11.9	4928	5	US-08-359-561-1	Sequence 1, Appl1
C 42	27.6	11.8	1230	5	US-08-890-719-3	Sequence 3, Appl1
C 43	27.6	11.8	11219	2	US-07-642-734C-1	Sequence 1, Appl1
C 44	27.6	11.8	11219	5	US-08-439-009A-1	Sequence 1, Appl1
C 45	27.4	11.7	3300	1	US-08-194-290-6	Sequence 6, Appl1

ALIGNMENTS

Pinda

RESULT 1
US-08-897-340-1

Patent No. 5955306

GENERAL INFORMATION:

; APPLICANT: Gimeno, Carlos J. and Errada, Patrick, R.
 ;
 ; TITLE OF INVENTION: Weight Control Pathway Genes and Uses
 ;
 ; TITLE OF INVENTION: Therefor

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSEE: LAHIVE & COCKFIELD, LLP
CITY: 38 State Street

STATE: 20 State Street
CITY: Boston
; ;

STATE: Massachusetts

COUNTRY: USA

; ZIP: 02109

; COMPUTER READ

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

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; OPENING SISIM: PC-DOS/MS-DOS
SOETMAPE: Patent In Release #1 0 Version #1 3E

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CURRENT APPLICATION DATA: COLIMANE: FACILITIN RELEASE W1:0,

APPLICATION NUMBER: US/08

FILING DATE: 7/21/07

CLASSIFICATION: 435-71

; PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 08/715,032
FILING DATE: 17 SEP 1996

FILED DATE: 17-SEP-1996

NAME: Silver, Jean M
ALL OTHER INFORMATION:
:

REGISTRATION NUMBER: 39,030

REFERENCE/DOCKET NUMBER: MNI-0050C

TELECOMMUNICATION INFORMATION:

TELEPHONE: (617) 227-7400 ;

TELEFAX: (617) 227-5941

```

; INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:

```

SEQUENCE CHARACTERISTICS:
LENGTH: 1386 base pairs

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: CDNA

US-08-897-340-1

Query match	100.0%;	Score 234;	DB 4;	Length 1386;
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Best Local Similarity 100.0%; Pred. No. 9.2e-68;
Matches 234; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

[illegible]

395 AGCTTGCACACATCCGACGATCAGCA

	Query Match	Similarity	Score	227.8	DB	4	Length	871
	Best Local	Similarity	97.9%	Pred.	No.	8,2e-66		
	Matches	229	Conservative	0	Mismatches	5	Indels	0
								Gaps
QY	1	acacagaccctgtcgtcgtacagaccatcgtcatcaagttygaagaatccgtgtcgc	60					
Db	335	ACACAGACCCTGTGCTGCGTAAACGACCTCATATGTCATATAATGTGAGAAATCCGTGCCG	394					
QY	61	agcttcagaccatcccgagacatcaacatctgtctgcagtgccctaccgcgcgggaactctt	120					
Db	395	AGCTTCAGACCATTCGCGAGATACCATTTGTTGTGGCAATGCCCTACCGCGGGAACATT	454					


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? CLASSIFICATION: 435
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US/08/361,705
? FILING DATE:
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: 07/966,040
? FILING DATE: 30-DEC-1992
? APPLICATION NUMBER: PCT/EP91/01361
? FILING DATE: 19-JUL-1991
? INFORMATION FOR SEQ ID NO: 11:
? SEQUENCE CHARACTERISTICS:
? LENGTH: 957 base pairs
? TYPE: nucleic acid
? STRANDEDNESS: single
? TOPOLOGY: linear
? MOLECULE TYPE: cDNA to mRNA
? ORIGINAL SOURCE:
? ORGANISM: Agkistrodon rhodostoma
? FEATURE:
? LOCATION: 210 to 911
? OTHER INFORMATION: The coding region shown in (2)(ix)(B)
? OTHER INFORMATION: codes for the protein of SEQ ID NO: 6
US-08-684-862-11

Query Match 15.3%; Score 35.8; DB 2; Length 957;
Best Local Similarity 55.1%; Pred. No. 0.0091;
Matches 70; Conservative 0; Mismatches 57; Indels 0; Gaps 0;

QY 13 tgcctcctaacgacctatgtctcatcaagttygaacgaatccgtgtccagtcgtgacacá 72
Db 454 TACTGGAACAAGACATATTGTGATCAAGCGTGAACCATCCTGTATGCAATAAGTGACACAA 513
QY 73 tcggagaataagacttcttcctcgacgtgcacctaccgcggggaactcttgecttgtttctg 132
Db 514 TGCGCCCTCTGACGTCCTGCCCTTCACGCCCTCCAGATGTGGGCTCATTTTGGCATATTATAG 573
QY 133 gctgagg 139
Db 574 GATGGGG 580

RESULT 8
US-08-684-862-12
? Sequence 12, Application US/08684862
? Patent No. 5759541
? GENERAL INFORMATION:
? APPLICANT: Bach, Alfred
? APPLICANT: Hillen, Heinz
? APPLICANT: Bialojan, Siegfried
? TITLE OF INVENTION: No. 5759541el Proteins, the Preparation and Use thereof
? NUMBER OF SEQUENCES: 14
? CORRESPONDENCE ADDRESS:
? ADDRESSEE: Keil & Weinkauff
? STREET: 1101 Connecticut Avenue
? CITY: Washington
? STATE: D.C.
? COUNTRY: USA
? ZIP: 20036
? COMPUTER READABLE FORM:
? MEDIUM TYPE: Diskette, 5.25 inch, 360 Kb storage
? COMPUTER: IBM AT-compatible, 80286 processor
? OPERATING SYSTEM: MS-DOS version 5.0
? SOFTWARE: Wordperfect version 5.1
? CURRENT APPLICATION DATA:
? APPLICATION NUMBER: US/08/684,862
? FILING DATE:
? CLASSIFICATION: 435
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US/08/361,705
? FILING DATE:
? PRIOR APPLICATION DATA:

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? APPLICATION NUMBER: 07/966,040
? FILING DATE: 30-DEC-1992
? APPLICATION NUMBER: PCT/EP91/01361
? FILING DATE: 19-JUL-1991
? INFORMATION FOR SEQ ID NO: 12:
? SEQUENCE CHARACTERISTICS:
? LENGTH: 840 base pairs
? TYPE: nucleic acid
? STRANDEDNESS: single
? TOPOLOGY: linear
?
JS-08-684-862-12

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Query Match	14.9%	Score 34.8;	DB 2;	Length 840;
Best Local Similarity	55.9%;	Pred. NO. 0.018;		
Matches 66; Conservative	0;	Mismatches 52;	Indels 0;	Gaps 0

Oy 22 acgagccatcattccatcaagtttgacgaatccggtcccgatctcgaacatccgagca 81
 376 AGGACATCATGTTGATCAGCGCTGAACAACTCTTAAACAACAGTAACACATCGCTCTC 435

RESULT 9
 US-08-467-155A-2
 Sequence 2, Application US/08467155A
 Patent No. 5736377
 GENERAL INFORMATION:
 APPLICANT: Band, Yumla
 TITLE OF INVENTION: NNS-1 POLYPEPTIDES, DNA, AND RELATED
 TITLE OF INVENTION: MOLECULES AND METHODS
 NUMBER OF SEQUENCES: 11
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Fish & Richardson P.C.
 STREET: 225 Franklin Street
 CITY: Boston
 STATE: MA
 COUNTRY: USA
 ZIP: 02110-2804
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/467,155A
 FILING DATE: 06-JUN-1995
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: Clark, Paul T.
 REGISTRATION NUMBER: 30,162
 REFERENCE/DOCKET NUMBER: 00398/100001
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 617/542-5070
 TELEFAX: 617/542-8906
 TELEX: 200134
 INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 1454 base pairs
 TYPE: nucleic acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: cDNA
 US-08-467-155A-2

Query Match	14.78;	Score 34.4;	DB 2;	Length 1454;
Best Local Similarity	52.98;	Pred. No. 0.031;		
Matches	74;	Mismatches	66;	Indels 0;
	Conservative			Gaps 0;

Accession	Sequence	Position
QY	22 acagccatcgtcatcaagtttgacgaatccggtgtccagttctgacacatccgagca	81
Db	488 ACGATCTATTTCTGCTAAGCTGGCCAGGCCCGTAGTCGCGGGGCCCGCGTCCGGGCC	547
QY	82 tcagatctgcttcgagtgccctaccgaggggaactcttgctctgtttctggtcgtggtc	141
Db	548 TGCACCTTCCTACCGCTGCTCTAGCCCGGAGACAGAGTGCACGTTGCTGCGGGCA	607
QY	142 tgcctgacgaagcagaatg	161
Db	608 CCACGGCGCGCCCGAGAGTG	627

RESULT 10
US-08-628-198-2
; Sequence 2, Application US/08628198
; Patent No. 5843694

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1  TITLE OF INVENTION:  NES-1 POLYPEPTIDES, DNA, AND RELATED
2  TITLE OF INVENTION:  MOLECULES AND METHODS
3  NUMBER OF SEQUENCES:  11
4  CORRESPONDENCE ADDRESS:
5  ADDRESSEE:  Fish & Richardson P.C.
6  STREET:  225 Franklin Street
7  CITY:  Boston
8  STATE:  MA
9  COUNTRY:  USA
10 ZIP:  02110-2804
11 COMPUTER READABLE FORM:
12 MEDIUM TYPE:  Floppy disk
13 COMPUTER:  IBM PC compatible
14 OPERATING SYSTEM:  PC-DOS/MS-DOS
15 SOFTWARE:  PatentIn Release #1.0, Version #1.30
16 CURRENT APPLICATION DATA:
17 APPLICATION NUMBER:  US/08/628,198
18 FILING DATE:
19 PRIOR APPLICATION DATA:
20 APPLICATION NUMBER:  08/467,155
21 FILING DATE:  06-JUN-1995
22 CLASSIFICATION:  514
23 ATTORNEY/AGENT INFORMATION:
24 NAME:  Clark, Paul T.
25 REGISTRATION NUMBER:  30,162
26 REFERENCE/DOCKET NUMBER:  00398/100002
27 TELECOMMUNICATION INFORMATION:
28 TELEPHONE:  617/542-5070
29 TELEFAX:  617/542-8906
30 TELEX:  200154
31 INFORMATION FOR SEQ. ID NO.:  2:
32 SEQUENCE CHARACTERISTICS:
33 LENGTH:  1454 base pairs
34 TYPE:  nucleic acid
35 STRANDEDNESS:  single
36 TOPOLOGY:  linear
37 MOLECULE TYPE:  cDNA
38  US-08-628-198-2

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Query Match	14.7%	Score 34.4;	DB 3;	Length 1454;
Best Local Similarity	52.9%	Pred. No. 0.031;		
Matches 74; Conservative	0;	Mismatches 66;	Indels 0;	Gaps 0;

[illegible]

COUNTRY: USA
ZIP: 20036
COMPUTER READABLE FORM:
MEDID TYPE: Diskette, 5.25 inch, 360 Kb storage
COMPUTER: IBM AT-compatible, 80286 processor
OPERATING SYSTEM: MS-DOS version 5.0
SOFTWARE: Wordperfect version 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/684,862
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/361,705
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/966,040
FILING DATE: 30-DEC-1992
APPLICATION NUMBER: PCT/EP91/01361
FILING DATE: 19-JUL-1991
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 1096 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA to mRNA
ORIGINAL SOURCE:
ORGANISM: Agkistrodon rhodostoma
FEATURE:
LOCATION: 144 to 841
OTHER INFORMATION: the coding region shown in (2)(ix)(B)
US-08-684-862-8

Query Match 14.2%; Score 33.2; DB 2; Length 1096;
Best Local Similarity 55.1%; Pred. No. 0.069; Mismatches 53; Indels 0; Gaps 0;
Matches 65; Conservative 0; Mismatches 53; Indels 0; Gaps 0;
QY 22 acgaaccatcatcgaatgagcaatccgtgctcgagtcgagtcgacacatccgagca 81
DB 403 AGACATCATGTTGATCAGCGTGAACAACTGTAAACAAGTGAACACATGCGTCC 462
QY 82 taagcatgcttcgcagtcgacctacgcgcggggaactcttgctcgttcttgctgag 139
DB 463 TCAGCTTGCTCCGCCCTCCCATGTGTGGCTCACTTGCCGTGTATGGGATGGGG 520
RESULT 14
US-08-557-146-1
Sequence 1, Application US/08557146
Patent No. 5834290
GENERAL INFORMATION:
APPLICANT: Egelrud, Torbjorn
APPLICANT: Hansson, Lennart
TITLE OF INVENTION: Recombinant Stratum Corneum Chymotryptic
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: White & Case, Patent Department
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2787
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/557,146
FILING DATE: 14-DEC-1995

CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Steiner, Richard J.
REGISTRATION NUMBER: 35,372
REFERENCE/DOCKET NUMBER: 1103326-181
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 819-8783
TELEFAX: (212) 354-8113
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 986 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
LOCATION: 25..786
FEATURE:
NAME/KEY: sig_peptide
LOCATION: 25..90
FEATURE:
NAME/KEY: mat_peptide
LOCATION: 112..783
US-08-557-146-1

Query Match 13.9%; Score 32.6; DB 3; Length 986;
Best Local Similarity 49.8%; Pred. No. 0.1;
Matches 114; Conservative 0; Mismatches 109; Indels 6; Gaps 1;
QY 1 acaaagaccctgctcgcgaacacctatgctcatcaagtcggagcaatccgyltcg 60
DB 335 ACTCCACACAGACCACCATGTTAATGACCTCATGTCTCGAAGCTCAATAGCCAGCCAGCC 394
QY 61 agctcgaacacatccgagcatcagatctgctcgagtcgacctacgcgcgggaactct 120
DB 395 TGTCATTCATGATGGAAGAAAGTCAGGCTGCCCTCCGCTCGAACCCTCGAACCACCT 454
QY 121 gctcgttcttgctgctgggtctgctgcga-----acggcagaatgctaccgyltcg 174
DB 455 GTACTGTCTCCGCCCTGCGGACACTACACAGCCAGATGTGACCTTCCCTTGACTCA 514
QY 175 agtcgctgaacgctgcgtggtgtctgaggaagtcgagtaagctcta 223
DB 515 TGTCGTGATGTCAAGCTCATCTCCGCCAGGACTGTCAGGAAGTTTA 563
RESULT 15
US-09-154-344-1
Sequence 1, Application US/09154344
Patent No. 5981256
GENERAL INFORMATION:
APPLICANT: Egelrud, Torbjorn
APPLICANT: Hansson, Lennart
TITLE OF INVENTION: Recombinant Stratum Corneum Chymotryptic
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: White & Case, Patent Department
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2787
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS

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? SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
? CURRENT APPLICATION DATA:
? APPLICATION NUMBER: US/09/154,344
? FILING DATE: 16-SEP-1998
? CLASSIFICATION:
? PRIOR APPLICATION DATA:
? APPLICATION NUMBER: US 08/557,146
? FILING DATE: 14-DEC-1995
? CLASSIFICATION:
? ATTORNEY/AGENT INFORMATION:
? NAME: Steiner, Richard J.
? REGISTRATION NUMBER: 35,372
? REFERENCE/DOCKET NUMBER: 1103326-181
? TELECOMMUNICATION INFORMATION:
? TELEPHONE: (212) 819-8783
? TELEFAX: (212) 354-8113
? INFORMATION FOR SEQ ID NO: 1:
? SEQUENCE CHARACTERISTICS:
? LENGTH: 986 base pairs
? TYPE: nucleic acid
? STRANDEDNESS: single
? TOPOLOGY: linear
? MOLECULE TYPE: cDNA
? HYPOTHEetical: NO
? ANTI-SENSE: NO
? ORIGINAL SOURCE:
? ORGANISM: Homo sapiens
? FEATURE:
? NAME/KEY: CDS
? LOCATION: 25..786
? FEATURE:
? NAME/KEY: s19-peptide
? LOCATION: 25..90
? FEATURE:
? NAME/KEY: mat-peptide
? LOCATION: 112..783
? US-09-154-344-1
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Query Match 13.98; Score 32.6; DB 4; Length 986;
Best Local Similarity 49.88; Pred. No. 0.1;
Matches 114; Conservative 0; Mismatches 109; Indels 6; Gaps 1;

QY 1 acaacagacccttgctgcctaacgacctatgctcatcaagtgtgacgaatccgtgtcgcg 60
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Db 335 ACTCCACACAGACCAAGCTTAATGACCTCATGCTGTGAAGCTCAATAGCCAGGCCAGGC 394

QY 61 agctgacacacatccgagatcagcatgtgttcgacagtgcctaccgcgagggaactctt 120
   || || || || || || || || || || || || || || || || || || || || ||
   395 TGTCAATCATGATGTGAAGAAAGTCAAGCTGCGCTCCGCTCGGAGAACCCCTGGAACCACT 454

QY 121 gctctgtttctgtgctgtgggtctgtgctgga-----acggcagaatgcttaacgtgtcgc 174
   || || || || || || || || || || || || || || || || || || || || ||
Db 455 GTACTGTCTCCGGCTGGGGCACTACACAGAGCCAGATGTGACCTTCCCTGACCTCA 514

QY 175 agtcggtgaagtgctgtgtgtgtgtcgtgagaggtcagtaagctcta 223
   || || || || || || || || || || || || || || || || || || || || ||
Db 515 TGTGCGTGATGTCAAGCTCATCTCCCGCCAGAGACTGACGAAGTTTA 563
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Search completed: December 15, 2000, 11:28:01
Job time: 44892 sec